Gamma oscillations in basal ganglia stem from the interplay between local inhibition and beta synchronization

Short title: Gamma band in basal ganglia stems from local inhibition and beta band

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Abstract

Basal ganglia activity fluctuations have primarily been studied in the context of beta (12-30 Hz) oscillations, a well-established neural marker for Parkinson's Disease (PD). Recent studies have also identified gamma (30-100 Hz) oscillations within the basal ganglia, suggesting it could serve as an alternative marker, but the underlying circuit mechanisms remain poorly understood. Here, through a spiking network model of the basal ganglia, we identified two distinct gamma oscillations: a high-frequency gamma rhythm within the globus pallidus (GPe-TI) and a slower gamma rhythm within D2 medium spiny neurons (MSNs), both stemming from self-inhibition. When we simulated dopamine depletion to mimic the effects of PD, the intensity of gamma oscillations in the GPe-TI was not affected, but their peak frequency increased due to phase-amplitude coupling with pathological beta oscillations. This suggests that the GPe-TI loop, while robust to dopamine depletion, becomes more synchronized with beta activity in the context of PD, leading to faster gamma rhythms. In contrast, gamma oscillations in D2 MSNs were not present in simulated healthy condition and only emerged

under dopamine-depleted pathological conditions. Moreover, both their intensity and peak frequency were strongly modulated by pathological beta activity.

Together, these findings highlight the complementary roles of self-inhibition and beta oscillations in shaping gamma activity within basal ganglia circuits. The GPe-TI loop primarily sustains high-frequency gamma rhythms, while low frequency gamma rhythms in D2 MSNs are strongly dependent on dopamine-depletion-related beta modulation. These results underscore the importance of network-wide interactions in PD, where pathological beta oscillations influence gamma activity. This study offers insights into the mechanisms of gamma oscillations in PD and highlights the potential of gamma activity, in both the prototypical and striatal loops, as a marker for disease progression and monitoring pathological dysfunction in PD.

Author Summary

Understanding the neural underpinnings of Parkinson's Disease (PD) is crucial for advancing treatments such as deep brain stimulation. While much research has focused on beta oscillations (12-30 Hz) in the basal ganglia as markers of Parkinsonian states, our study explores the role of gamma oscillations (30-100 Hz), which have been observed experimentally but are less understood. Using a computational model, we identified two distinct types of gamma rhythms: a high-frequency oscillation in the globus pallidus and a slower oscillation in D2 medium spiny neurons.

Our findings reveal that in simulated healthy conditions, gamma oscillations originate within the globus pallidus. However, under conditions mimicking PD, gamma activity in D2 MSNs is also present. Both rhythms stem from local inhibition and are shaped by pathological beta oscillations in both frequency and spectral power.

These results show how gamma oscillations originate in the basal ganglia and how they strongly interact with pathological beta synchronization related to PD. Our study highlights the potential of gamma activity as a valuable marker for guiding deep brain stimulation and understanding the underlying dysfunctions in PD.

Introduction

The basal ganglia (BG) are a group of interconnected subcortical nuclei in the brain that play a pivotal role in regulating motor control, learning, emotions, and decision-making. The main structures that comprise the basal ganglia are the striatum (STR), globus pallidus (GP), subthalamic nucleus (STN), and substantia nigra (SN) (1). The striatum contains fast-spiking interneurons (FSNs) and medium-spiny neurons (MSNs), which express D1- and D2-type dopamine receptors, while the GP is subdivided into the external (GPe) and internal (GPi) segments. These nuclei are interconnected in a complex network that helps modulate voluntary movement and other cognitive functions (2). Disruptions in the functioning of these structures, particularly in the context of neurodegenerative diseases, can lead to profound motor and behavioral impairments. Parkinson's disease (PD) is the second most common neurodegenerative disorder, affecting approximately 0.3% of the global population. The disease is primarily characterized by motor symptoms such as akinesia (3), bradykinesia (4), resting tremor (5), and gait disturbances (6). PD is caused by the progressive degeneration of dopaminergic neurons in the substantia nigra pars compacta (7), leading to a depletion of dopamine in the striatum and disruptions in basal ganglia signaling. This depletion specifically affects D1- and D2-receptor-expressing neurons, which alters the dynamic balance of activity within the basal ganglia network. As PD progresses, these disruptions manifest as motor impairments and are often managed through pharmacological treatments or surgical interventions. One such intervention is deep brain stimulation (DBS), a surgical approach that delivers electrical pulses to specific basal ganglia nuclei, typically the STN or GPi, to alleviate motor symptoms (8,9). However, conventional DBS delivers continuous stimulation without accounting for fluctuations in symptom severity or variations in the patient's daily activities. This limitation has spurred the development of adaptive DBS (aDBS), a more advanced form of stimulation that adjusts in real-time based on the patient's physiological signals (10–12). aDBS requires continuous monitoring of brain activity to modulate stimulation parameters in response to dynamic changes in motor function. A critical challenge for aDBS is identifying electrophysiological biomarkers that reliably correlate with symptoms, reflect pathophysiological mechanisms, and support precise real-time modulation of electrical stimulation.

Among the electrophysiological changes observed in PD (13–17), oscillatory activity in the basal ganglia has been the subject of intense study. Of these oscillations, beta-band activity (12–30 Hz) has garnered the most attention. Beta oscillations are prominently observed in the STN and GPi of PD patients, and their amplitude

correlates with the severity of motor impairment (18–21). Moreover, beta oscillations were found in both rats (22) and monkeys (23,24) models of PD, confirming the key role of these oscillations in the disease. These oscillations are now widely considered as biomarkers for DBS, particularly for adjusting aDBS (11). However, beta-band oscillations are influenced by various factors, including DBS stimulation, medication, and task performance, which can complicate their use in real-time modulation (25–27). This inherent variability has led to growing interest in gamma oscillations (30–100 Hz) as an alternative, more stable biomarker for adaptive DBS (28–30).

Gamma oscillations have emerged as a significant biomarker across a range of pathological conditions (31), including PD. These oscillations have been observed in the STN and other basal ganglia structures in both PD patients (32–36) and animal models (37–40). They are modulated by movement (41,42), associated with dyskinetic states (43), and linked to sleep disturbances commonly observed in PD (44). Unlike beta oscillations, gamma rhythms show a stronger robustness to external influences such as DBS stimulation and medication, making them a promising candidate for real-time monitoring in aDBS systems (29,30). Moreover, recent studies have demonstrated that gamma oscillations are tightly correlated with motor dysfunction (33), suggesting their potential role in PD symptomatology.

Despite the growing interest in gamma oscillations, the precise mechanisms underlying their generation and modulation in the basal ganglia network are not fully understood. Computational modeling, which has been used extensively to study basal ganglia dynamics, serves as a valuable tool to address these open problems (45–47). While some computational models have explored gamma rhythms in the BG, most have focused on beta oscillations and treated gamma as a secondary epiphenomenon. For example, Hodgkin-Huxley-based models have shown that gamma oscillations (~80 Hz) can emerge in FSNs and GPe, with STN-DBS restoring these rhythms following dopamine depletion (48). Other studies have highlighted specific BG loops, such as the GPe-STN network, as key players in generating gamma rhythms in the 35–50 Hz range (49,50). Additionally, spiking models have linked gamma activity in the STN and GPe to motor control tasks (51) or variations in MSN connectivity (52). However, the role of gamma oscillations within the broader BG circuitry remains poorly understood.

In this study, we extend a previously established computational model (53) to explore the dynamics of gamma oscillations in the basal ganglia. We focus on the role of local inhibition in driving gamma rhythms in the

striatum and GPe and examine how dopamine depletion, synaptic parameters, and external input strengths influence these oscillations. Our work aims to provide new insights into the mechanisms of gamma oscillations in PD and propose a theoretical framework for their potential use as biomarkers in adaptive DBS systems.

Results

Gamma oscillations in a computational model of basal ganglia

We investigated dopamine-depletion-driven oscillations in the basal ganglia (BG) firing activity starting from a previously validated spiking network model (53) (see Methods and Fig 1A). Coherently with previous findings (53), beta-range oscillations (12–30 Hz) were well-defined across all BG nuclei under healthy conditions (dopamine depletion level, $D_d = 0.75$; black lines in Fig 1B). Dopamine depletion enhanced the power of these beta oscillations in all nuclei ($D_d = 1$; orange lines in Fig 1B and S2A Fig), reproducing the emergence of pathological beta activity observed in the PD condition.

In contrast to the widespread beta oscillations, activity in the high gamma range (~100 Hz) was observed only in the GPe-TI nucleus under healthy conditions (Fig 1B). Under dopamine-depleted states, gamma oscillations persisted in GPe-TI and newly emerged in the D2 population (Fig 1B). Interestingly, gamma oscillations in GPe-TI showed little to no change in power with increasing dopamine depletion (Fig 1C; Spearman $\rho = 0.19$, $p = 0.36$), while their peak frequency exhibited a non-monotonic trend (S2B Fig, bottom). In contrast, gamma activity in the D2 population was highly sensitive to dopamine depletion: gamma power increased significantly with greater depletion levels (Fig 1C; Spearman $\rho = 0.99$, $p << 0.001$), with significant oscillations emerging only for $D_d > 0.87$ and showing an increase in peak frequency at higher depletion levels (S2C Fig, bottom).

These findings highlight two key insights. First, gamma oscillations in the D2 population are sensitive to dopamine dynamics, suggesting they may reflect pathological changes associated with Parkinson's disease progression. Second, GPe-TI and D2 populations exhibit differential sensitivity to dopamine depletion, with gamma oscillations in GPe-TI being more resilient, while those in D2 increase in intensity and frequency with dopamine deficiency. Furthermore, our results reveal a distinct relationship between beta- and gamma-range oscillatory dynamics during dopamine depletion. In the following sections, we will explore the mechanistic origins of these oscillations and discuss their implications for understanding the emergence and regulation of abnormal rhythmic activity in the basal ganglia during PD progression.

Globus pallidus self-inhibition drives gamma oscillations features

To explore the role of specific synaptic pathways in generating GPe-TI gamma oscillations within our basal ganglia model, we systematically silenced each connection individually and analyzed the resulting effects on GPe-TI gamma power (see Methods). Specifically, we calculated the ratio of GPe-TI gamma power with each connection silenced to the GPe-TI gamma power in the intact full-BG model (Fig 2A). Our results demonstrated that gamma oscillations were still present after the removal of most connections (Fig 2A), with one notable exception: GPe-TI self-inhibition. Disrupting this self-inhibitory pathway led to a near-complete loss of gamma oscillatory activity (Fig 2B), underscoring GPe-TI self-inhibition as a necessary mechanism for gamma rhythmogenesis within this nucleus. Interestingly, the removal of GPe-TI self-inhibition was also associated with a marked increase in beta oscillations power (Fig 2B).

To further investigate the role of GPe-TI self-inhibition in regulating gamma oscillations, we systematically reduced the strength of self-inhibitory feedback and assessed its effect on GPe-TI gamma power. To maintain a consistent average input to GPe-TI despite the reduced self-inhibition, we introduced an auxiliary Poisson inhibitory input (see Methods and Fig 2C, where it is referred to as "GPe-TI aux."). However, this compensatory input was insufficient to fully stabilize the average firing rate of the GPe-TI population, which consequently increased as the strength of self-inhibition decreased (S3A Fig, left).

Our findings revealed that progressively weakening self-inhibition synaptic strength $\alpha_{Gpe\text{-}TI}$, led to a gradual reduction in gamma activity (Fig 2D, Spearman $\rho = 1.00$, $p \ll 0.001$) and peak frequency (S3A Fig, right and Spearman $\rho = 0.86$, $p \ll 0.001$). When self-inhibition was fully silenced, gamma activity disappeared, declining to levels of power below statistical significance (Fig 2D). Notably, we also observed a concurrent increase in beta synchronization as self-inhibition strength decreased (Fig 2D, Spearman $\rho = -0.96$, $p \ll 0.001$) suggesting an inverse relationship between GPe-TI gamma oscillations (or, equivalently, self-inhibition strength) and beta hypersynchrony.

We next examined how other characteristics of GPe-TI self-inhibition (specifically, synaptic decay time and delay) impacted gamma activity. We found that the increase of both delay and decay times of GPe-TI self-inhibition synapses reduced the gamma peak frequency (Fig 2E top-left and top-middle. Delay: Spearman = -0.99, $p \ll 0.001$. Decay time: Spearman $\rho = -0.99$, $p \ll 0.001$), though their effects on gamma power differed: increasing synaptic delay boosted gamma power (Fig 2E middle-left, Spearman $\rho = 0.98$, $p \ll 0.001$), whereas

increasing decay time reduced it (Fig 2E middle, Spearman $\rho = -0.98$, $p \ll 0.001$). Notably, the reduction in gamma power associated with increased decay time was accompanied by a decrease in the average firing rate of GPe-TI (Fig 2E bottom-middle; Spearman $\rho = -1.00$, $p \ll 0.001$). In contrast, the increase in gamma power due to prolonged synaptic delays occurred without a corresponding increase in GPe-TI firing rate (Fig 2E bottom-left, Spearman $\rho = 0.54$, $p = 0.005$), suggesting that this enhancement reflected improved synaptic efficacy rather than increased firing activity alone.

We also observed distinct effects of decay time and delay on beta oscillations. Interestingly, we found beta oscillations power to closely reflect the average GPe-TI firing activity. GPe-TI self-inhibition delay had little to no effect on both beta power (Fig 2E middle-left, Spearman $\rho = 0.53$, $p = 0.007$) and firing rate. Conversely, increased decay times significantly reduced both beta and GPe-TI firing (Fig 2E middle, Beta power: Spearman $= -0.99$, $p \ll 0.001$). Taken together, these findings suggest that while beta oscillations follow the average firing rate of GPe-TI, gamma oscillations are influenced not only by firing rate but also by local resonance properties within GPe-TI that are not reflected in beta oscillations.

Finally, we examined the effect of excitatory drive on GPe-TI gamma activity. As expected, increasing the excitatory drive to GPe-TI raised its average firing rate (Fig 2E bottom-right Spearman $\rho = 1.00$, $p \ll 0.001$), which in turn elevated both beta and gamma power (Fig 2E middle-right. Beta power: Spearman $= 0.99$, $p \ll 0.001$. Gamma power: Spearman $\rho = 0.99$, $p \ll 0.001$), as well as the gamma peak frequency (Fig 2E top-right, Spearman $\rho = 0.99$, $p \ll 0.001$). This finding underscores a link between GPe-TI firing activity and gamma oscillatory dynamics, with excitatory input modulating gamma features primarily through its effect on firing rate. However, as shown earlier, gamma oscillations also depend on intrinsic local dynamics within GPe-TI, suggesting that gamma power reflects a combination of both firing rate and the synaptic properties of the recurrent inhibition.

Gamma oscillations in the globus pallidus are modulated by beta activity

In the previous section, we demonstrated that GPe-TI self-inhibition was necessary for the presence of gamma oscillations in this nucleus. Here, we assessed whether this self-inhibitory loop alone was also sufficient for gamma rhythmogenesis. To this end, we analyzed an isolated GPe-TI population, configured with the same parameters as the original network but disconnected from all other basal ganglia nuclei. Accordingly, the inputs

from other nuclei were replaced with auxiliary constant-rate Poissonian inputs that matched the firing activity of their respective original projections (in Fig 3A, indicated with “GPe-TA aux.”, “D2 aux.” and “STN aux.”). We found that, under healthy conditions, the isolated GPe-TI firing spectrum closely resembled that of the full BG network within the gamma range (Fig 3B, top; R^2=0.86 for frequencies between 50 and 150 Hz, see Methods). However, the isolated GPe-TI was unable to display beta oscillations, which are known to arise from the interplay among different BG nuclei (53). Under PD conditions, the behavior of the isolated GPe-TI diverged significantly from that of the full BG network (Fig 3B, bottom; R^2 = 0.34). As expected, the pathological beta hypersynchrony, which arises from interactions across BG nuclei, was entirely absent in the isolated GPe-TI. Furthermore, gamma activity in the isolated GPe-TI was notably diminished, showing reductions in both spectral power and peak frequency relative to the full BG network (Fig 3B). This suggests that, while self-inhibition was sufficient to sustain gamma oscillations under healthy conditions, additional inter-nuclei interactions are critical for maintaining both beta and gamma dynamics under pathological dopamine-depleted states.

To investigate this discrepancy further, we analyzed the effect of dopamine depletion on gamma power and peak frequency in both the isolated and full BG scenarios (Fig 3C). For low dopamine depletion levels (D_d < 0.87), the gamma power and peak frequency of the isolated and full BG GPe-TI populations remained similar, as indicated by a high goodness-of-fit between the firing spectra of the two scenarios (S4A Fig, left). However, beyond this threshold (D_d > 0.87), their behaviors began to diverge (S4A Fig, left): while gamma power and peak frequency increased in the full BG network, they progressively declined in the isolated GPe-TI. Interestingly, near this critical threshold ($D_d \approx 0.87$), we observed that pathological beta power in the full BG network began to dominate and override gamma activity. This finding suggests that the observed differences between the isolated and full BG cases arise from the increasingly intense beta oscillations in the full network. To support this hypothesis, we compared the GPe-TI firing rates across dopamine depletion levels (S5A Fig). The average firing rates of the isolated and fully connected GPe-TI were comparable, decreasing progressively with dopamine depletion (S5A Fig, left). However, the standard deviation of the firing rates showed a marked divergence between the two scenarios (S5A Fig, right). In the isolated GPe-TI, the standard deviation decreased alongside the average firing rate, which led to a corresponding reduction in gamma power and peak frequency. This pattern reflected a direct relationship between firing dynamics and gamma oscillations in the absence of

223 broader network interactions. In contrast, the full BG network exhibited a different behavior. While the average
224 firing rate decreased with dopamine depletion, the presence of beta oscillations caused the standard deviation
225 of GPe-TI firing rates to increase with higher levels of dopamine depletion (S5A Fig, right). These beta
226 oscillations, driven by interactions across BG nuclei, induced significant fluctuations in GPe-TI firing rates,
227 particularly at higher dopamine depletion levels (Fig 3D). Such fluctuations transiently elevated GPe-TI firing
228 rates, leading to an increase in gamma power and peak frequency, an effect absent in the isolated GPe-TI
229 scenario where beta oscillations were not present.
230 To quantitatively explore the relationship between beta and gamma oscillations in the full BG model, we
231 analyzed the beta-triggered scalogram of GPe-TI firing rates under PD conditions (Fig 3E). This analysis
232 revealed that both gamma spectral power and peak frequency were strongly influenced by the phase of beta
233 oscillations. Specifically, gamma activity reached its highest power and frequency at a beta phase of
234 approximately 0, corresponding to the peak of GPe-TI firing observed in the full BG model (Fig 3D). Crucially,
235 these fluctuations were absent in the firing of the isolated GPe-TI due to the absence of beta oscillations (Fig
236 3D and S5A Fig, right).
237 This coupling between gamma oscillations and beta phases explains the differences in gamma power and peak
238 frequency observed in the PSDs of the isolated and full BG cases. The PSD provides an average power
239 estimation over the entire time axis, disproportionately emphasizing the higher-frequency components of
240 gamma oscillations, as these coincide with periods of elevated power driven by beta synchronization.
241 Consequently, the PSD of GPe-TI in the full BG model exhibited gamma oscillations of greater power and
242 higher peak frequency compared to the isolated case (Fig 3B, C), driven by the transient beta-triggered
243 increases in firing rates. To test whether the isolated GPe-TI could replicate these dynamics, we analyzed the
244 relationship between its average firing rate and the resulting gamma power and frequency (Fig 3E). As
245 expected, we found a positive correlation between these factors (firing rate and gamma peak frequency:
246 Spearman $\rho = 1.00$, $p \ll 0.001$; firing rate and gamma power: Spearman $\rho = 1.00$, $p \ll 0.001$; gamma peak
247 frequency and power: Spearman $\rho = 1.00$, $p \ll 0.001$), reinforcing the hypothesis that the enhanced gamma
248 activity in the full BG model is attributable to transient increases of GPe-TI firing rates driven by beta
249 oscillations.

250 Nonetheless, adjusting the average firing rate of the isolated GPe-TI revealed its limitations in fully replicating
251 the spectral properties of the full BG model (S4A Fig, right). Even with an optimized GPe-TI firing rate that
252 maximized the goodness-of-fit (maximum R^2=0.80 for a firing rate of 45.22 spike/s), the match remained
253 incomplete. This discrepancy highlights the critical role of beta-driven fluctuations and broader network
254 interactions in shaping gamma oscillations (S4A Fig, right; maximum R2=0.80 for a firing rate of isolated
255 GPe-TI equal to 45.22 spike/s).
256 Overall, these results underscore that GPe-TI self-inhibition is both necessary and sufficient for the emergence
257 of gamma oscillations under both healthy and pathological conditions. However, in pathological conditions,
258 beta-driven fluctuations (arising from network interactions beyond self-inhibition) play a critical role in
259 modulating gamma activity, a dynamic that cannot be fully replicated by simply adjusting the nucleus's average
260 firing rate.
261
262 **D2 self-inhibition shapes gamma oscillations features**
263 Following the same approach used for the GPe-TI nucleus, we systematically silenced each synaptic pathway
264 individually in the full-BG model and analyzed its impact on D2 gamma power. Specifically, we calculated
265 the ratio of D2 gamma power with each connection silenced to the gamma power in the intact full-BG model
266 (Fig 4A). Since gamma oscillations in D2 neurons were absent in dopamine depletion levels corresponding to
267 the healthy condition, this analysis was conducted under the PD condition. Our findings revealed that
268 disrupting D2 self-inhibition nearly abolished gamma oscillatory activity (Fig 4A and Fig 4B). This disruption
269 was accompanied by a significant increase in beta oscillation power (Fig 4B), suggesting that D2 self-inhibition
270 plays a critical role in sustaining gamma oscillations in the PD state while simultaneously suppressing beta
271 activity.
272 To further investigate the role of D2 self-inhibition in regulating gamma oscillations, we systematically
273 reduced the strength of self-inhibitory feedback and assessed its effect on D2 gamma power. To maintain a
274 consistent average input to D2 despite the reduced self-inhibition, we introduced a compensatory Poisson input
275 to offset the diminished inhibitory drive (see Methods and Fig 4C, where it is indicated with "D2 aux."). Unlike
276 what was observed for GPe-TI, this compensatory input was sufficient to maintain a stable D2 average firing
277 across self-inhibition synaptic strengths (S3B Fig, left).

Our findings revealed that gradually reducing the strength of D2 self-inhibition (α_{D2}) led to a progressive decrease in gamma oscillation power (Fig 4D). This decline continued until gamma oscillations completely disappeared when the D2-D2 synaptic strength was reduced to 40% of the original reference value (Fig 4D; Spearman $\rho = 0.96$, $p \ll 0.001$). At this point, gamma power fell below the threshold for statistical significance. A similar trend was observed for the D2 gamma peak frequency (S3B Fig, right; Spearman $\rho = 0.80$, $p < 0.001$). In contrast, beta oscillations in D2 neurons exhibited an opposing trend, with power increasing as the D2-D2 synaptic strength decreased (Fig 4B; Spearman $\rho = -0.97$, $p \ll 0.001$). This inverse relationship between gamma and beta oscillations in D2 neurons closely mirrored the results observed in GPe-TI.

Beyond synaptic strength, we investigated how other characteristics of D2 self-inhibition (specifically, synaptic decay time and delay) impacted gamma activity. Increasing the synaptic decay time of D2-D2 connections caused a slight decrease in gamma power (Fig 4E, middle; Spearman = -0.88, $p \ll 0.001$), gamma frequency (Fig 4E, top-middle; Spearman $\rho = -0.95$, $p \ll 0.001$), and the average firing rate of D2 neurons (Fig 4E, bottom-middle; Spearman $\rho = -0.98$, $p \ll 0.001$). This aligns with the notion that longer decay times reduce the overall excitability of the nucleus by prolonging inhibitory input, leading to a reduction in firing activity and, consequently, in gamma power and frequency.

In contrast, altering the synaptic delay of D2-D2 connections had minimal effect on the average firing rates of D2 neurons (Fig 4E, bottom-left; Spearman $\rho = 0.11$, $p = 0.59$). However, synaptic delay exerted distinct and opposing effects on gamma oscillations: gamma power increased (Fig 4E, middle-left; Spearman $\rho = 0.98$, $p \ll 0.001$), while gamma frequency decreased (Fig 4E, top-left; Spearman $\rho = -0.99$, $p \ll 0.001$). These findings suggest that synaptic delay primarily modulates the timing and efficiency of self-inhibitory interactions within D2 neurons, thereby affecting gamma oscillations more directly and independently of firing rate changes. This pattern mirrors the behavior observed in the GPe-TI, where changes in synaptic decay times affected firing rates and gamma oscillations in tandem, while changes in synaptic delay acted selectively on gamma features.

Finally, we assessed the impact of modulating external excitatory drive to D2 neurons on gamma oscillations, a manipulation that mimics the effects of dopamine depletion and models the transition from the healthy to the Parkinsonian condition. As expected, increasing the excitatory drive to D2 neurons elevated their average firing rate (Fig 4E, bottom-right; Spearman $\rho = 1.00$, $p \ll 0.001$), which was accompanied by simultaneous increases

306 in both gamma and beta power (Fig 4E, middle-right; gamma power: Spearman $\rho = 1.00$, $p \ll 0.001$; beta
307 power: Spearman $\rho = 0.96$, $p \ll 0.001$) as well as in gamma peak frequency (Fig 4E, top-right; Spearman $\rho =$
308 0.90, $p \ll 0.001$). Interestingly, within the range of D2 external drive levels corresponding to healthy to PD
309 transition, both beta and gamma oscillations emerged as hallmarks of shifting network dynamics. Notably, beta
310 oscillations appeared earlier during the transition from healthy to pathological state, while gamma oscillations
311 became prominent only at later stages. These findings highlight the potential of striatal D2 gamma oscillation
312 power and frequency as sensitive markers for tracking dopamine depletion and the progression between
313 healthy and PD conditions.
314
315 **Gamma oscillations in D2 are modulated by beta activity**
316 In the previous section, we demonstrated that D2 self-inhibition is a necessary condition for the emergence of
317 gamma activity within the nucleus. Analogously to what was done for GPe-TI, we next investigated whether
318 the D2 self-inhibitory loop alone was also sufficient for gamma rhythmogenesis. To address this, we analyzed
319 an isolated D2 population configured with the same parameters as the original network but disconnected from
320 all other basal ganglia nuclei. In this setup, inputs from other nuclei were replaced with auxiliary constant-rate
321 Poisson inputs calibrated to match the firing activity of their corresponding original projections (Fig 5A,
322 indicated with "FSN aux.", "D1 aux." and "GPe-TA aux.").
323 Under healthy conditions, we found that the spectral patterns of D2 firing activity in the isolated nucleus and
324 the full BG network were poorly aligned (Fig 5B, R^2=0.05), as neither scenario exhibited gamma activity.
325 However, under PD conditions, gamma oscillations emerged in both the isolated D2 population and the full
326 BG model. Despite this, the consistency between the two spectral patterns remained weak (Fig 5B, R^2=0.49),
327 with the isolated D2 population displaying reduced gamma power and central frequency compared to the full
328 BG scenario. Furthermore, as expected, the pathological beta hypersynchrony observed in the full BG D2 was
329 absent in the isolated case.
330 To further investigate this discrepancy, we analyzed the effect of dopamine depletion on gamma power in both
331 the isolated D2 and full BG scenarios (Fig 5C). In the full BG model, increasing dopamine depletion led to a
332 concurrent rise in both beta and gamma power. Notably, gamma oscillations in this scenario crossed the
333 threshold of statistical significance at intermediate levels of dopamine depletion, approximately halfway

through healthy and PD conditions. A similar dependence on dopamine depletion was observed in the isolated D2 nucleus, but the magnitude of gamma oscillations was consistently lower than in the full BG network. As a result, gamma power in the isolated D2 remained below the significance threshold until very high levels of dopamine depletion were reached. Accordingly, increasing dopamine depletion level led to only a mild improvement in the goodness-of-fit between the firing spectra of the scenarios (S4B Fig, left).

To elucidate the mechanisms underlying the reduced gamma power in the isolated D2 nucleus, we explored the relationship between beta and gamma oscillations in the full BG model. This approach was motivated by the observation that beta oscillations played a key role in enhancing gamma activity in the GPe-TI (Fig 3). To quantitatively examine this relationship, we computed the beta-triggered scalogram of D2 firing rates under PD conditions (Fig 5E). This analysis was restricted to PD states because D2 gamma oscillations were absent in healthy conditions.

Our analysis revealed that both gamma spectral power and peak frequency were strongly modulated by the phase of beta oscillations. Specifically, gamma activity in the D2 population transitioned rhythmically between non-significant and significant power values as a function of the beta cycle phase. Consistent with our findings in the GPe-TI, D2 gamma power and frequency peaked at a beta phase corresponding to the maximum D2 firing rate observed in the full BG model (Fig 5D). In contrast, these beta-driven fluctuations were absent in the isolated D2 population due to the lack of beta oscillations (Fig 5D and S5B Fig, right).

These results indicate that the average firing rate of D2 neurons is insufficient to sustain gamma rhythmogenesis when considered in isolation. Instead, beta oscillations in the full BG model transiently elevate D2 firing rates, driving the nucleus into a firing regime where its self-inhibitory dynamics have sufficient energy to sustain gamma rhythmogenesis. This beta-mediated mechanism highlights the critical role of inter-nuclear interactions in modulating gamma activity under pathological conditions.

To determine whether the lack of gamma activity in the isolated scenario could be exclusively resolved by modulating D2 average firing activity, we analyzed the relationship between gamma power and the average firing rate of the isolated D2 population. As expected, increasing the average D2 firing rate led to a corresponding increase in gamma activity within the nucleus. Specifically, we found that an increase of 2.8 sp./s was required for the isolated D2 to match the gamma power observed in the full BG network. However, adjusting the firing rate of D2 alone was insufficient to fully replicate the spectral characteristics of gamma

activity in the full BG circuitry (S4B Fig, right; maximum R^2=0.70 for a firing rate of isolated D2 equal to 4.33 spike/s).

Overall, these findings demonstrate that D2 self-inhibition is necessary but not sufficient for gamma rhythmogenesis, under both healthy and pathological conditions. The emergence and modulation of gamma oscillations critically depend on beta-driven fluctuations, which arise from inter-nuclear interactions within the full basal ganglia circuitry. This underscores the importance of network-wide dynamics in shaping gamma activity and highlights the limitations of isolated nuclei in fully replicating the spectral properties of the full system.

Discussion

Using a computational spiking model of the basal ganglia (53), we identified two distinct types of gamma activity, originating from separate nuclei (Fig 1): D2 receptor-type medium spiny neurons in the striatum and the prototypical population of the globus pallidus. These two types of gamma oscillations exhibited different characteristics. D2 gamma oscillations occurred exclusively under pathological conditions, were confined to lower frequencies, and showed a marked increase in power with dopamine depletion. By contrast, GPe-TI gamma oscillations were observed consistently across healthy and pathological states, with their power remaining unaffected by dopamine depletion.

Despite these distinctions, our findings revealed a unifying principle underlying both rhythms: their generation relies critically on recurrent inhibition within their respective nuclei. The removal of inhibitory connections in the model completely abolished gamma oscillations in both cases (Fig 2 and Fig 4), underscoring the centrality of this mechanism. Furthermore, altering the synaptic time scales of self-inhibition modulated the spectral power and frequencies of these oscillations, further supporting the pivotal role of local inhibitory circuits in gamma activity patterns.

While recurrent inhibition proved necessary for gamma rhythmogenesis in both nuclei, it alone could not fully account for the observed gamma dynamics. In the GPe-TI, self-inhibition was sufficient to generate gamma oscillations (Fig 3); however, the specific frequency and power of these oscillations observed in the complete BG model required interactions with beta oscillations. In contrast, gamma rhythmogenesis in D2 MSNs

required not only recurrent inhibition but also phase-amplitude coupling with beta oscillations, highlighting a more complex interplay of mechanisms in this population (Fig 5).

The spiking network model presented in this work is largely based on a previously presented model by our group (53). In the previous work, we focused on the relationship between dopamine depletion and beta oscillations, hence gamma oscillations were not investigated. The version of the network presented in (53), however, displayed high-frequency oscillations exceeding the range of the currently available experimental results (S2 Table). The novel version of the network described in the present work accounts instead for both beta and gamma oscillations in basal ganglia. For a detailed comparison of the modifications introduced in the network see the dedicated subsection "Comparison with previous version of the network" in the Methods and S3 Table.

Our findings are consistent with prior computational models that identify the GPe as the primary source of basal ganglia gamma oscillations (48–51). These models similarly reported the limited dependence of GPe gamma oscillations on dopamine depletion levels, despite variations in the oscillation frequencies across studies. Among these, only (49) explicitly incorporated the two subpopulations of the GPe, aligning with our identification of the prototypical population as the driver of gamma activity.

The literature on striatal gamma oscillations presents a more heterogeneous picture. For example, findings from a model focusing exclusively on striatal populations (52) align with our results, identifying D2 MSNs as the primary source of gamma oscillations. In contrast, a different model (48) attributes gamma activity in the striatum to fast-spiking interneurons rather than D2 MSNs. This discrepancy may stem from fundamental differences in model design, particularly the absence of explicit modeling of D1 and D2 MSN subpopulations in the latter study. Such differences highlight the role of neuronal subtypes in determining complex dynamics of gamma oscillations in the striatum.

Experimental evidence from local field potential recordings in rats further supports our findings, with gamma oscillations observed in both the striatum and in the GPe (37–40,54). GPe gamma oscillations have been suggested to arise locally (40), consistently with our results. Similarly, striatal gamma oscillations have been found to correlate with beta power (38) and were shown to intensify with increased dopamine depletion (55), though this latter study reported lower gamma frequencies than those predicted by our model, highlighting an area for further investigation.

Although calibrated using rat data, our model might provide insights into gamma oscillations observed in the STN of human patients (32,33,56). While our spiking model does not directly simulate LFPs, which reflect synaptic inputs rather than spiking activity, it can inform hypotheses about their origins. Under the hypothesis that LFPs in basal ganglia nuclei arise from synaptic inputs, as within the cortex (57), STN gamma activity could reflect the gamma-patterned firing of its inputs, particularly from the GPe-TI, which projects directly to the STN. Our findings support this possibility, suggesting that GPe-TI gamma activity may underlie gamma oscillations experimentally observed in STN LFPs. In contrast, striatal gamma oscillations are unlikely to influence STN activity due to the lack of direct connections between these two nuclei (2).

A notable outcome of this study is the anti-correlation between pathological beta oscillations and D2 and GPe-TI self-inhibition. Interestingly, experimental recordings in 6-OHDA-lesioned rats have demonstrated an increase in GPe-TI self-inhibition under Parkinsonian conditions (58). This increase does not appear to result directly from changes in dopamine receptor activity within GPe-TI self-inhibition, suggesting that this enhanced synaptic strength reflects a form of adaptive homeostasis aimed at counterbalancing pathological beta oscillations.

The coupling between beta-phase and gamma-amplitude in the context of Parkinson's Disease has been experimentally observed between the subthalamic nucleus and the motor cortex (59), within the motor cortex (60,61) and within the subthalamic nucleus (59,62), in both rodent models and human patients. Here, we characterized the mechanisms behind beta-gamma phase-amplitude coupling within the basal ganglia, demonstrating how beta-induced fluctuations of GPe-TI and D2 firing activities can influence gamma patterns. This provides a plausible explanation for the experimentally observed coupling between these two bands, shedding light on how beta oscillations can influence gamma-generating nuclei through intrinsic network interactions.

Despite the strong alignment of our results with both experimental and computational studies, our computational framework has certain limitations. Experimental evidence indicates that calcium (63) and hyperpolarization-activated cyclic nucleotide-gated (64) currents are essential for the bursting activity of the STN. These ionic processes were not included in our model due to the simplicity inherent in the choice of neuronal models adopted in the computational framework. Specifically, we employed integrate-and-fire point neuron models, which prioritize computational efficiency over the detailed representation of complex ionic

dynamics. This design choice enabled large-scale network simulations but inevitably excluded certain biophysical intricacies that could further refine model's accuracy in future works.

Another key assumption in our study concerns the inputs provided to all populations within the BG circuitry, which were modeled as uncorrelated Poisson processes. While this approach simplified the complexity of network inputs, it does not account for the broader context in which the BG functions as part of an interconnected network involving the cortex and thalamus (1). Evidence suggests that beta and gamma oscillations in the BG may be, at least in part, inherited from cortical activity (65–67). To achieve a more comprehensive understanding of these rhythms, future studies will then need to incorporate more biologically plausible models of cortical and thalamic inputs to capture their contributions to BG dynamics. Although such integrations were beyond the scope of this study, they represent an important direction for future work.

Overall, our findings demonstrate that the basal ganglia are capable of intrinsically generating gamma oscillations, independent of cortical or thalamic inputs, highlighting the self-sustained nature of these rhythms. Importantly, our results do not preclude the possibility that gamma oscillations may also be influenced by, or inherited from, extra-BG sources such as cortical or thalamic inputs. Instead, our modeling supports the hypothesis that the BG can independently generate and sustain gamma rhythms through intrinsic mechanisms.

Our modeling of dopamine depletion introduces another limitation. In this study, dopamine depletion was represented as an increased firing rate in the D2 populations, reflecting the heightened activity of the indirect pathway in PD (45). While effective in reproducing key features of pathological states, this simplification does not account for other critical factors, such as changes in synaptic strengths, plasticity mechanisms, or network-level adaptations that occur in Parkinson’s disease (68). By focusing on the most direct consequence of dopamine depletion (i.e., the imbalance between the direct and indirect pathways of the basal ganglia (46)) we were able to capture significant changes in both beta and gamma oscillatory activity. However, future iterations of the model could benefit from incorporating these additional factors to provide a more complete representation of the disease state.

In conclusion, our work offers the first comprehensive explanation of the origin of gamma oscillations in the basal ganglia and their modulation under pathological conditions. By elucidating the mechanisms underlying gamma rhythms and their interaction with beta oscillations, this study advances our understanding of basal

ganglia dynamics. Furthermore, it highlights the potential of gamma oscillations as biomarkers for adaptive deep brain stimulation, paving the way for improved therapeutic strategies for Parkinson's disease.

Material and Methods

Basal ganglia model

The basal ganglia model used in this work was adapted from a previous one implemented to study the origins of beta oscillations (Fig 1A and (53)).

The model architecture comprised the striatum, the external globus pallidus and the subthalamic nucleus. The striatum was modeled with three neural populations: D1 receptor-type MSNs, D2 receptor-type MSNs and FSNs. The external globus pallidus was instead divided into two populations, according to experimental recordings in the pallidus of rats (69,70): the prototypic (GPe-TI) and arkypallidal populations (GPe-TA). The subthalamic nucleus was instead the only excitatory population of the model. The population numerosities of the model are reported in Table 1.

The connectivities among populations were random with fixed probabilities based on previous works (53,68). Table 2 summarizes the connection properties within the BG model.

All populations in the model received external excitatory Poissonian inputs, simulating afferent drive from other brain structures. These inputs were calibrated to adjust each population's firing rate within a range consistent with experimental observations (see S1 Table and S1 Fig). The external input rates and synaptic strength are reported in Table 3.

Simulations were performed through the neural simulator ANNarchy (71). The integration method adopted was the 4th-order Runge-Kutta (72), with stepsize 0.1 ms. Each simulation of five seconds was repeated 5 times (except for the raster plots in Fig 3 and Fig 5, panel D). In all cases, we discarded an additional 1 second of simulation to avoid the effects of the initial conditions.

Population	FSN	D1	D2	GPe-TA	GPe-TI	STN
N° of neurons	420	6000	6000	264	780	408

Table 1. **Population numerosities of the six populations of the BG model.**

Source	Target	Probability	Delay [ms]	Type	Synaptic weight [nS]
D1	D1	0.0607	1.7	Inh.	0.12
D1	D2	0.0140	1.7	Inh.	0.30
D2	D1	0.0653	1.7	Inh.	0.36
D2	D2	0.0840	1.7	Inh.	0.25
D2	GPe-TI	0.0833	7.0	Inh.	1.28
FSN	D1	0.0381	1.7	Inh.	6.60
FSN	FSN	0.0238	1.0	Inh.	0.50
FSN	D2	0.0262	1.7	Inh.	4.80
GPe-TI	GPe-TI	0.0321	1.8	Inh.	1.1
GPe-TI	GPe-TA	0.0321	1.0	Inh.	0.35
GPe-TI	FSN	0.0128	7.0	Inh.	1.60
GPe-TI	STN	0.0385	1.0	Inh.	0.08
GPe-TA	D1	0.0379	7.0	Inh.	0.35
GPe-TA	D2	0.0379	7.0	Inh.	0.61
GPe-TA	FSN	0.0379	7.0	Inh.	1.85
GPe-TA	GPe-TA	0.0189	1.0	Inh.	0.35
GPe-TA	GPe-TI	0.0189	1.0	Inh.	1.20
STN	GPe-TA	0.0735	2.0	Exc.	0.13
STN	GPe-TI	0.0735	2.0	Exc.	0.42

Table 2. Connection probabilities, synaptic delays and weights for all the connections within the BG model.

Target nucleus	ν_{ext} [spike/s]	Synaptic weight [nS]
D1	1276	0.45
D2	900*	0.45
FSN	944	0.50
GPe-TI	600	0.25
GPe-TA	170	0.15
STN	500	0.25

Table 3. Parameters of the external excitatory input to each nucleus of the model. "*" indicates the value for healthy condition. For PD conditions, the excitatory input to D2 was set to 1200 spike/s (see "Modeling of dopamine depletion").

Neuronal models

All the neurons in this work were modelled as adaptive conductance-based integrate-and-fire (73,74). Following the modeling approach described in (53,68), our model resulted from the combination of two previously implemented networks: the striatal model of (75,76) and the pallidus-subthalamic circuit model of (77). Accordingly, it also inherited their choice of using two different point neuron models: quadratic integrate-and-fire for striatal populations and exponential integrate-and-fire for pallidus-subthalamic nuclei.

Specifically, striatal populations (D1, D2 and FSN) were modeled as adaptive quadratic integrate-and-fire (78):

$$C_m \frac{dV(t)}{dt} = k(V(t) - E_L)(V(t) - V_{th}) - g_{ex}(t)(V - E_{ex}) - g_{in}(t)(V(t) - E_{in}) - w(t) + I_e \qquad (1)$$

where k scales the voltage-dependent term, V is the membrane potential, E_L is the leak reversal potential, V_{th} is the threshold potential for spike initiation, g_{ex} and g_{in} are the excitatory and inhibitory conductances, E_{ex} and E_{in} are their respective reversal potentials, w represents the adaptation variable, and I_e is an external current.

Pallidal (GPe-TI and GPe-TA) and subthalamic neurons were considered instead as adaptive exponential integrate-and-fire (78):

$$C_m \frac{dV(t)}{dt} = -g_L(V(t) - E_L) - g_{ex}(t)(V(t) - E_{ex}) - g_{in}(t)(V(t) - E_{in}) + g_L \Delta T e^{\left(\frac{V(t) - V_{th}}{\Delta T}\right)} - w(t) + I_e \quad (2)$$

where C_m is the membrane capacitance, ΔT the slope factor determining the sharpness of the exponential activation near the threshold, and the exponential term represents the nonlinear exponential term modeling action potential generation.

For both models, if $V > V_{peak}$ a spike is emitted and the membrane potential is reset to V_{reset}.

In all cases but for FSN, the dynamic of the adaptive variable was described by:

$$\tau_w \frac{dw(t)}{dt} = -w(t) + a(V(t) - E_L) + b\tau_w \sum_i \delta(t - t_i) \qquad (3)$$

where τ_w is the adaptation time constant, a scales the subthreshold adaptation, b determines the strength of spike-triggered adaptation, and the sum accounts for discrete spike times of the neuron t_i.

For FSN, according to the modeling of (75), the dynamics for the adaptive variable was driven by:

$$\tau_w \frac{dw(t)}{dt} = b\tau_w \sum_{t_i} \delta(t - t_i) + \begin{cases} -w(t) + a(V(t) - V_b)^3, & if\ V(t) < V_b \\ -w(t), & otherwise \end{cases} \qquad (4)$$

where the parameter V_b and the two regimes allow for a non-linear transition from silent to spiking dynamics. The dynamics of excitatory and inhibitory synapses are described by the time evolution of conductances in each post-synaptic neuron, as represented by the following equations:

$$\begin{cases} \tau_{ex}\dfrac{dg_{ex}(\mathrm{t})}{dt} = -g_{ex}(t) + \tau_{ex} \displaystyle\sum_{exc.\ neur.\ j} g_j \sum_i \delta\left(t - t_i^j - t_r^j\right) \\ \tau_{in}\dfrac{dg_{in}(t)}{dt} = -g_{in}(t) + \tau_{in} \displaystyle\sum_{inh.\ neur.\ j} g_j \sum_i \delta\left(t - t_i^j - t_r^j\right) \end{cases} \tag{5}$$

where τ_{ex} and τ_{in} represent the decay times for excitatory and inhibitory synaptic conductances of the postsynaptic neurons, respectively, and g_{ex} and g_{in} the corresponding synaptic conductances. g_j denotes the synaptic weight from the presynaptic neuron j, t^j_i refers to the spike time i of the presynaptic neuron j, t_r^j is the synaptic delay associated with the synapses originating from neuron j and δ is the Dirac delta function. The outer summations account for the contributions of all presynaptic neurons connected to the postsynaptic neuron under consideration, while the inner summations consider all spike events of the presynaptic neuron j.

All the neuronal parameters are equal to those of (53), are reported in Table 4. The synaptic weights and delays can be found instead in Table 2.

Parameter	Unit	D1	D2	FSN	GPe-TI	GPe-TA	STN
C_m	pF	15.2	15.2	80.0	40.0	60.0	60.0
E_L	mV	-78.2	-80.0	-80.0	-55.1	-55.1	-80.2
E_{ex}	mV	0.0	0.0	0.0	0.0	0.0	0.0
E_{in}	mV	-74	-74	-74	-65	-65	-84
τ_{ex}	ms	12.0	12.0	12.0	10.0	10.0	4.0
τ_{in}	ms	10.0	10.0	10.0	7.0	5.5	8.0
V_{th}	mV	-29.7	-29.7	-50.0	-54.7	-54.7	-64.0
I_e	pA	0.0	0.0	0.0	12.0	1.0	5.0
t_{ref}	ms	0.0	0.0	0.0	0.0	0.0	0.0
V_{reset}	mV	-60	-60	-60	-60	-60	-70
a	nS	-20	-20	0.025	2.5	2.5	0.0
b	pA	67.0	91.0	0.0	70.0	105.0	0.05
τ_w	ms	100.0	100.0	5.0	20.0	20.0	333.0
V_{peak}	mV	40.0	40.0	25.0	15.0	15.0	15.0
ΔT	mV	-	-	-	1.7	2.55	16.2
g_L	nS	-	-	-	1.0	1.0	10.0
k	nS/mV	1.0	1.0	1.0	-	-	-
V_b	mV	-	-	-55	-		-

Table 4. **Parameters for the neuronal equation of each population of the BG model.**

Modeling of dopamine depletion

Our work aimed to investigate gamma oscillation patterns in healthy and Parkinsonian conditions. Parkinson's disease has profound effects on striatal dynamics, which are particularly sensitive to the degeneration of dopaminergic neurons in the substantia nigra pars compacta (79). Experimental studies have shown that dopamine depletion alters the excitability of projection neurons in a receptor-specific manner (80). Specifically, neurons expressing D1 and D2 dopamine receptors exhibit opposite responses to dopaminergic inputs (81,82). In Parkinsonian conditions, this differential response results in an overall increase in the firing activity of D2 neurons, accompanied by a corresponding decrease in the firing activity of D1 neurons (83,84).

In our computational model, we directly implemented this increase in D2 neuron activity by modulating the rate of the external excitatory input, adopting the same assumption of (53). Specifically, the transition from healthy to PD was modeled with a dopamine depletion (D_d) parameter according to the rule:

$$\nu_{ext}(D_d) = D_d \nu_{ext}^0 \quad (6)$$

where the reference value ν_{ext}^{0} is equal to 1200 spike/s. The parameter D_d was then varied from 0.75 (representing healthy conditions) to 1 (for simulating the Parkinsonian condition).

This adjustment increased the D2 firing rate by elevating the dopamine depletion parameter D_d, effectively capturing the dynamics observed in PD without explicitly modeling dopamine signaling. Moreover, given the inhibitory nature of D2 neurons and their projections onto D1 neurons, enhancing D2 activity led to a corresponding reduction in D1 firing without the need for direct modulation of D1 external input.

Comparison with the model's previous version

In this work, we used a newer version of the model compared to our previously developed one (53). The reason for this update was that, although the original model was capable of generating gamma oscillations (an aspect not explicitly reported in (53), see Discussion), it exhibited excessively high frequencies of pallidal gamma activity, reaching approximately 130 Hz. While gamma oscillations were not the primary focus of the previous study, adjusting their frequency was essential for our investigation, as experimentally observed gamma activity in GPe-TI falls within a lower frequency range (S2 Table). The revised parameter set effectively reduced the gamma oscillation frequency while preserving the beta rhythms and maintaining the original firing dynamics of the model.

The parameter adjustments primarily targeted the GPe-TI-GPe-TI and D2-D2 connections, as these pathways and their parameters significantly influence gamma oscillations (see sections "Globus pallidus self-inhibition drives gamma oscillations features" and "D2 self-inhibition shapes gamma oscillations features" in Results).

For GPe-TI self-inhibition the synaptic delay was set to 1.8 ms, and the decay time of all the connections to GPe-TI was increased from 5.5 ms to 7 ms. These changes reduced the central frequency of GPe-TI gamma oscillations by approximately 30 Hz while keeping the parameter values consistent with other computational models (Synaptic delay of GPe-TI to GPe-TI: ~[2-4] ms (85–87); decay time of GPe-TI: ~[5-8] ms (87)). The synaptic weights for GPe-TI self-inhibition were reduced from 1.2×10^{-3} to 1.1×10^{-3}, and the external input rate to GPe-TI neurons was decreased from 1530 spikes/s to 600 spikes/s. These adjustments were necessary to counterbalance the increase in gamma power caused by the lower synaptic delay (Figure 2E, left column).

For the D2-D2 connections, the synaptic weight was increased from 0.2×10^{-3} to 0.25×10^{-3}, and the external input rate to D2 neurons was scaled by a factor of 1.11, enhancing the excitatory drive. Additionally, the

external input to D1 neurons was increased from 1127 spikes/s to 1280 spikes/s. These changes were essential to restore the firing activity of D2 and D1 neurons and to recreate the gamma power of D2 gamma oscillations reported in (53), which had been affected due to modifications in the GPe-TI pathway.

A summary of the changes made in our work as compared to (53) is reported in S3 Table.

Spectral analysis

All spectral analyses were performed on the population firing rates, defined as the number of spikes produced by a population within a 1 ms time window divided by the product between the size of the population and the window length. Specifically, for the spectral analysis we computed the power spectral densities (PSDs) using Welch's method (88), with 1 second of window length, 500 ms of overlap, and Hamming window.

If present, beta harmonics (see Fig 1B) were removed with a spectral interpolation technique (89), to avoid bias in the computation of gamma spectral power and frequency.

Although the power spectral densities from simulated data exhibit reduced complexity compared to experimentally measured ones, they still reflect spectral properties influenced by synaptic filtering, which introduced spurious trends unrelated to oscillatory dynamics (90). Since these trends could cause an overestimation in the computations of the spectral power, they were removed by fitting this aperiodic component and subtracting it. We chose to model the aperiodic component as a power law function of the frequency:

$$Ap(f;\alpha,\beta) = 10^{\beta}\frac{1}{f^{\alpha}} \tag{7}$$

where α is the exponent of the power law and β an offset value. The fitting procedure was performed with the FOOOF method (91), which was used to identify the aperiodic components of the PSDs.

Once the beta harmonics and the aperiodic components were removed from the PSDs, we computed the oscillation frequency in both beta and gamma bands. This was estimated with the "centre of gravity" method (92):

$$f_{peak} = \frac{1}{\int_{f_1}^{f_2} PSD(f)df}\int_{f_1}^{f_2} fPSD(f)df \tag{8}$$

where we chose f_1=10 Hz and f_2=30 Hz for the beta band, f_1=50 Hz and f_2=150 Hz for GPe-TI gamma and f_1=40 Hz and f_2=120 Hz for D2 gamma. We adopted different ranges for the two gamma oscillations because GPe-TI and D2 gamma oscillations presented peaks at different frequencies (Fig 1B), and selecting specific ranges improved the accuracy of the peak identification.

We then used the computed central frequencies to choose the band on which to compute the mean spectral power:

$$\langle PSD \rangle = \frac{1}{f_2 - f_1} \int_{f_1}^{f_2} PSD(f) df \tag{9}$$

Specifically, we considered the beta and gamma bands respectively as $\left[f_{peak}^{\beta} - 5, f_{peak}^{\beta} + 5\right] Hz$ and $\left[f_{peak}^{\gamma} - 20, f_{peak}^{\gamma} + 20\right] Hz$. The use of bands centred around the peak frequencies was required to avoid changes in the spectral power due to the shift in frequency of the corresponding peak.

We also implemented a threshold to determine whether the power spectral densities displayed significant spectral power in the considered band. Specifically, the spectral power of the firing activity in the considered range was compared with that of fictitious Poissonian processes with the same average firing rate. To this purpose, 100 equivalent activities with a constant mean rate $\nu(t) = \nu_0$ were simulated. We generated the number of spiking neurons n_j in each time bin j of the synthetic process from a binomial distribution:

$$p(n_j; \nu_0, N) = B(n_j; \nu_0, N) = \binom{N}{n_j} \nu_0^{n^j} (1 - \nu_0)^{N - n_j} \tag{10}$$

where N is the number of neurons in the considered population and ν_0 the probability of spiking each neuron, computed as the mean number of spikes of the population divided by the corresponding numerosity. We then estimated the threshold with the spectral power of the synthetic population in the chosen band, averaged across the 100 realisations of the process. If the mean spectral power of the original population exceeded this threshold, a significant oscillation was considered present in the activity. The thresholds were reported with dashed lines in all the figures representing analyses on the spectral powers.

Identification of most relevant gamma oscillations

To identify which of the populations of the model presented significant gamma oscillations (Sections "Gamma oscillations in a computational model of basal ganglia" and S1 Fig), we computed the gamma spectral power of activity of each nucleus, considering the gamma range as $[50,120]$ Hz. Unlike all the others, for this analysis we adopted a fixed range for the gamma band to avoid any assumption about potential gamma oscillations. Furthermore, we selected a broad range of frequencies to capture any source of gamma activity and to account for the wide distribution of the observed gamma peaks. We then compared the spectral powers with those of the Poissonian processes in the same band and considered significant oscillations only those that exceeded this threshold (see section "Spectral analysis" in Materials and Methods). With this approach, we found that two populations displayed gamma oscillations in their activities: GPe-TI and D2 (See Fig 1B and 1C). Consequently, all the other analyses of this work were performed exclusively on these two populations.

Identification of necessary projections for gamma oscillations

We aimed to determine whether gamma rhythms in the BG model depended on specific synaptic connections. To investigate this, we systematically disconnected each projection in the model and analyzed how gamma oscillations in the two nuclei were affected (Fig 2B and Fig 4B). Synaptic disconnection was implemented by setting the corresponding synaptic weights to zero. However, this removal altered the input received by the target population, potentially affecting not only the spectral pattern of its firing dynamics but also its average level. To compensate for the missing input while isolating the role of the connection in generating gamma oscillations, we introduced an auxiliary Poissonian population. This population matched the size and average firing rate of the original source population and provided a white noise input to the target nucleus through a projection with the same characteristics as the removed one. For instance, if the disconnected projection was from D2 to GPe-TI, an auxiliary Poissonian population with the same size and average firing rate as D2 was created to send input to GPe-TI through a projection with synaptic properties identical to the original "D2 to GPe-TI" connection. By maintaining a constant average input to the target nucleus, this approach ensured that overall firing dynamics remain stable within the BG model. However, it effectively eliminated any temporal structure originally present due to the disconnected synaptic projections. This allowed us to determine whether the specific temporal patterns of firing activity originating from a given projection contributed to the generation

of gamma oscillations or whether their influence was merely due to the average input level of the source nucleus.

To compare the spectral power in GPe-TI (Fig 2B) and D2 (Fig 4B) with and without each connection, we then calculated the ratio Rγ of gamma power with each connection silenced to the gamma power in the intact model. In this way, we were able to identify those projections whose presence was the most relevant to gamma rhythmogenesis.

Investigation of the effects of recurrent synaptic parameters on oscillations

The disconnection of the model projections described in the previous section allowed for the identification of self-inhibition as necessary condition for gamma oscillations (Fig 2B and Fig 4B). To further characterize the role of these connections (i.e., D2 to D2 and GPe-TI to GPe-TI), we investigated the effects of their synaptic parameters on gamma oscillations (sections "Globus pallidus self-inhibition drives gamma oscillations features" and "D2 self-inhibition shapes gamma oscillations features" in Results).

First, we analyzed the effects of reducing self-inhibitory strength in D2 and GPe-TI on the spectral power of both beta and gamma oscillations. This was achieved by scaling down the synaptic weights of recurrent inhibitory connections by a multiplicative factor $\alpha \in [0,1]$. Since this reduction weakened self-inhibition and consequently altered the firing dynamics of the nucleus, we introduced an auxiliary Poissonian population (labeled 'aux.' in Fig 2C and Fig 4C) to compensate for the loss of inhibitory input. This auxiliary population matched the original nucleus in terms of neuron count, synaptic parameters, and average firing rate. It provided external input through a projection with synaptic weights scaled by $1-\alpha$ relative to the original self-inhibition, ensuring that the overall level of inhibitory input remained unchanged. However, because the auxiliary population generated Poisson-distributed activity, this substitution eliminated the temporal structure inherent to the original self-inhibitory projections. This approach allowed us to isolate the role of recurrent inhibition in shaping gamma oscillations while preserving the nucleus's mean input.

Next, we examined the effects of altering the synaptic delay and decay time of self-inhibition in GPe-TI and D2 on their firing rates, gamma spectral power, and gamma peak frequencies (Fig 2E and Fig 4E, left and center columns). We also studied how the change in the average rate of the external excitatory drive to GPe-TI and D2 affected these quantities (Fig 2E and Fig 4E, right column).

All relationships between the alteration of synaptic parameters and the observed quantities were evaluated using the Spearman coefficient, with the corresponding p-value computed via a bootstrap permutation test with 1,000 permutations. The investigations were performed in the pathological case (D_d=1), because in healthy conditions only GPe-TI gamma oscillations were present.

Isolation of gamma-generating nuclei

In addition to investigating whether self-inhibitory connections were necessary for gamma oscillations, we examined whether they were also sufficient to sustain gamma activity in GPe-TI and D2 (see sections "Gamma oscillations in the globus pallidus are modulated by beta activity" and "Gamma oscillations in D2 are modulated by beta activity" in Results). To this end, we created isolated versions of GPe-TI and D2, generating populations ("isolated GPe-TI" and "isolated D2" in Fig 3A and Fig 5A, respectively) that retained the same parameters and neuron count as in the full model. These populations were disconnected from all other nuclei while preserving their recurrent inhibitory connections and external excitatory Poissonian drive.

A key challenge of this isolation was to maintain firing rates comparable to those in the full model despite these removed inputs. To achieve this, we replaced the disconnected projections with inputs from auxiliary Poissonian populations. Similarly to what was detailed in previous sections, these auxiliary populations generated Poisson-distributed synaptic inputs with the same average firing rate and synaptic properties as those of the original source populations they replaced. This approach ensured that the isolated nuclei received the same average input as in the full model while eliminating any temporal (and thus spectral) structure present in the original projections. For instance, in the full BG model, GPe-TI receives inputs from GPe-TA, D2, and STN. In the isolated GPe-TI condition, these inputs were replaced by Poissonian versions of these nuclei, labeled GPe-TA aux., D2 aux., and STN aux. (Fig 3A). This substitution preserved the average level of synaptic inputs while preventing any spectral influence present in the original network activity, allowing us to assess whether self-inhibition alone was sufficient to generate gamma oscillations.

We then compared the spectra of the isolated populations with the corresponding original ones, in both healthy and PD (Fig 3B and Fig 5B). Specifically, we computed the coefficient of determination R^2 between the two power spectral densities (original and isolated), considering only the frequencies between 50 Hz and 150 Hz

to avoid bias induced by the presence of beta oscillations. The PSDs considered for this analysis were averaged across five simulations, to limit noise variability.

The isolated populations were also used to compute the relationship between the firing rate of the population and the spectral power and peak frequency of gamma oscillations (Fig 3F and Fig 5F).

Beta-induced modulations on gamma oscillations

Analysis of the population activity in nuclei exhibiting gamma oscillations suggested a potential coupling between beta and gamma rhythms (Fig 3D and Fig 5D). To explore this interaction, we conducted a time-frequency decomposition of the firing activities by calculating their beta-averaged scalogram (Fig 3E and 5E). Specifically, we first applied a Morlet wavelet transform (5 cycles, 200 frequency steps between 60 and 150 Hz) to the firing activity of each nucleus to obtain its time-resolved spectral power. Next, we extracted the beta phase by applying a Hilbert transform to the bandpass-filtered signal (Butterworth filter, 5th order, [10–30] Hz). The beta phase was then discretized into 100 bins, and the scalogram was averaged within each phase bin. This procedure allowed us to quantify how the power of different frequencies varied across the beta cycle, providing insight into the phase-dependent modulation of gamma oscillations.

To distinguish genuine gamma oscillations from background activity, we compared these amplitudes to those obtained from a Poisson process with the same average firing rate as the original nucleus. Specifically, we computed the wavelet transform of the firing rate of a Poissonian spiking process with the same duration and mean rate as the original data. The resulting gamma-band wavelet amplitudes were then used as a significance threshold: only frequency-phase bins where the original amplitude exceeded the Poisson-derived values were considered indicative of significant oscillatory activity.

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Author contribution

F.F.: Conceptualization, Formal Analysis, Software, Visualization, Writing – Original Draft Preparation; M.A.: Conceptualization, Software, Supervision, Writing – Original Draft Preparation; N.M.: Conceptualization, Supervision, Visualization, Writing – Original Draft Preparation; E.C.: Conceptualization, Project Administration, Supervision, Writing – Review & Editing; A.M.: Conceptualization, Funding Acquisition, Project Administration, Supervision, Writing – Original Draft Preparation, Writing – Review & Editing.

Data and code availability statement

The codes necessary to reproduce the data that support the findings of this study are openly available at https://github.com/CNELab/Gamma-Oscillations-Basal-Ganglia.

Figures

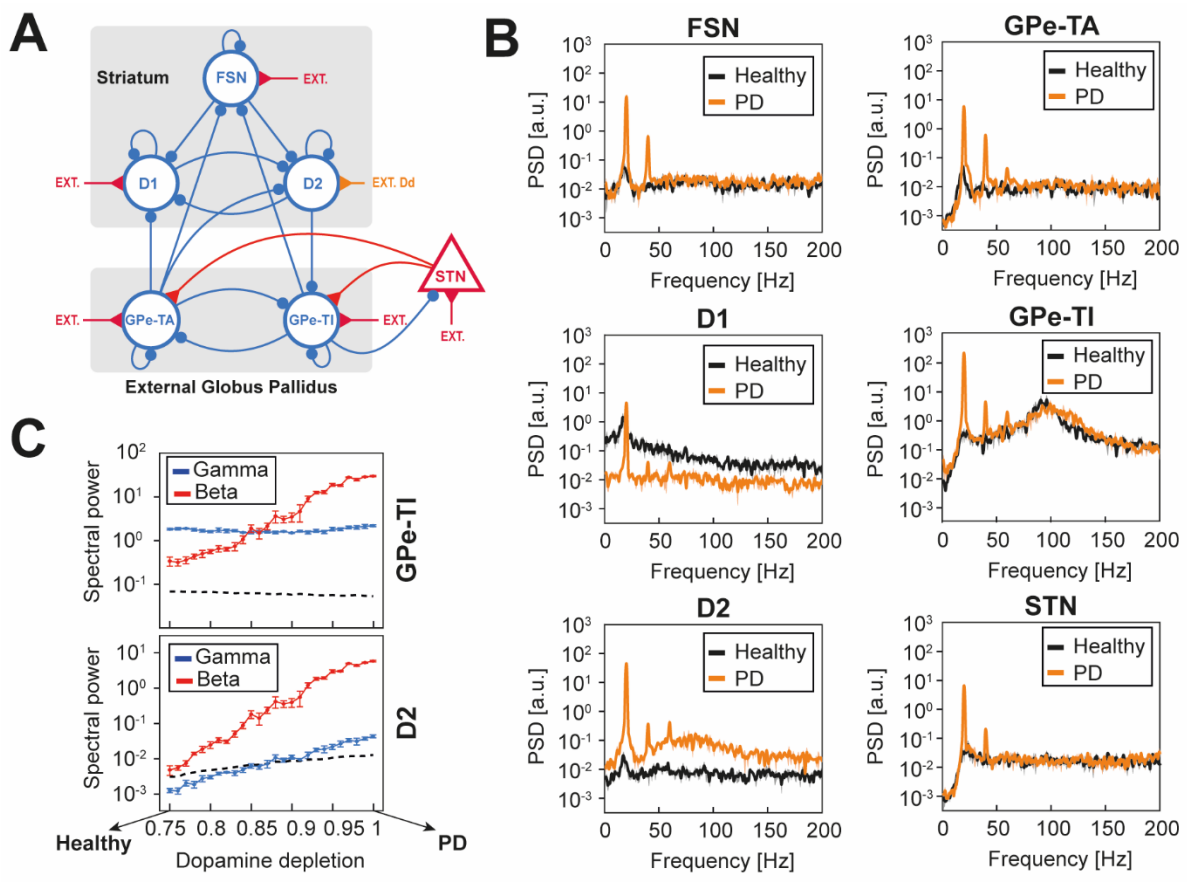

Fig 1. Gamma oscillations in the globus pallidus and the striatum of a basal ganglia spiking model. (A) Architecture of the BG model. FSN, D1 and D2: striatal spiking interneurons, and medium spiny neurons with D1 and D2 dopamine receptors; GPe-TA and GPe-TI: arkypallidal and prototypic populations of the globus pallidus pars externa; STN: subthalamic nucleus; ext: external excitatory Poissonian input. Red/blue arrows indicate excitatory/inhibitory projections. Dopamine depletion (D_d, orange external drive) was modeled by adjusting the excitatory Poisson input to D2 neurons. (B) Power spectral densities (PSDs) of each BG nucleus activity under healthy (D_d= 0.75, black) and Parkinson's disease conditions (D_d= 1, orange). PSDs are reported as mean and standard deviation across five simulations of 5 seconds. (C) Spectral powers of GPe-TI (top) and D2 (bottom) firing activity within beta (red) and gamma (blue) ranges across dopamine depletion levels. Data are reported as mean and standard deviation across five simulations. The threshold for statistically

significant oscillatory power (dashed black line) was determined considering the power spectrum of an equivalent Poisson process with matched firing rate.

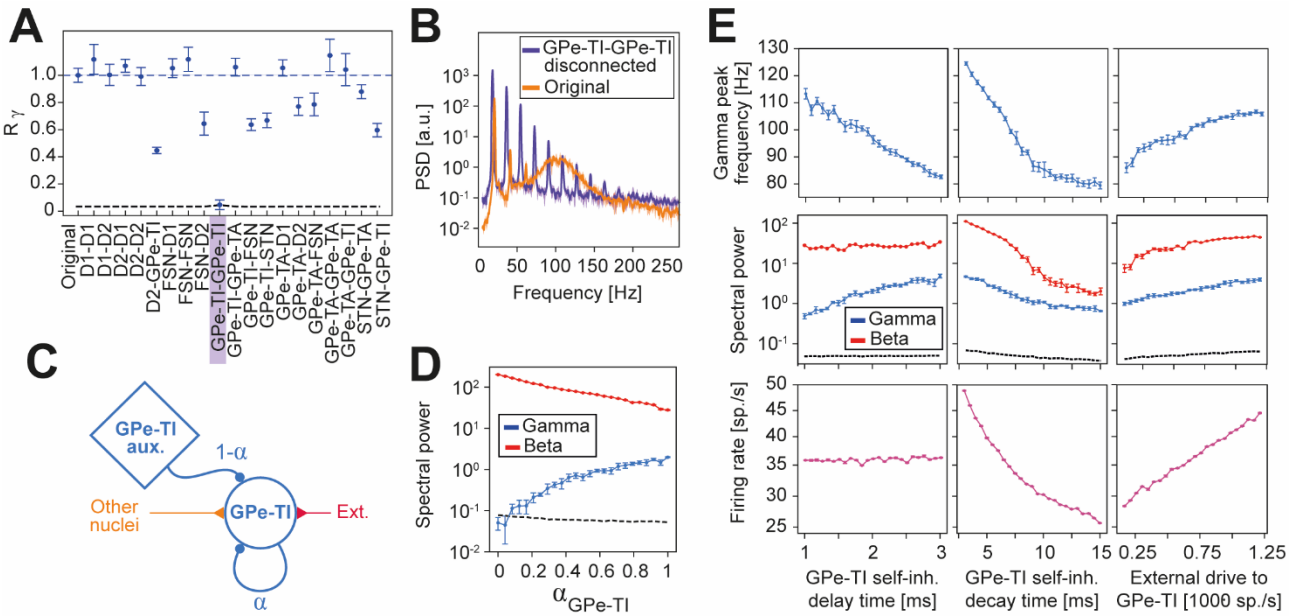

Fig 2. Self-inhibition parameters and external drive modulate GPe-TI oscillations. (A) Ratio R_γ of GPe-TI gamma power with each synaptic connection silenced compared to the GPe-TI gamma power in the intact full-BG model. The spectral power ratio of GPe-TI activity in the gamma band is shown with ('original') and without each synaptic connection in the full-BG model, with the specific connection removed indicated on the x-axis. For each removed connection, the threshold (black dashed line) for significant oscillations is computed as the ratio between the power of equivalent poisson processes and the gamma power in the fully connected model. The blue dashed line indicates the spectral power ratio for the fully connected case. (B) Power spectral densities (PSDs) of GPe-TI with (orange) and without (purple) self-inhibitory connection, in Parkinsonian condition (D_d=1). (C) Scheme illustrating the procedure for reducing GPe-TI self-inhibition: the weights of self-inhibitory connections were progressively decreased by a multiplicative factor α Є [0,1], and input from an auxiliary Poissonian population (GPe-TI aux.) was added to compensate for the reduction in self-inhibitory input. The afferent activity from other BG nuclei and external inputs were left unchanged. (D) GPe-TI spectral power of beta (red) and gamma (blue) bands across self-inhibition synaptic strength α. The threshold for statistically significant oscillatory power (dashed black line) was determined considering the power spectrum of an equivalent Poisson process with matched firing rate. (E) Gamma peak frequency (top row), beta (red)

and gamma (blue) spectral powers (middle row), and GPe-TI firing rate (bottom row) across GPe-TI self-inhibition delay (left column), GPe-TI self-inhibition decay time (middle column), and the strength of external input to GPe-TI (right column). In the middle row, the threshold for significant oscillations was indicated with a black dashed line. All data was reported as mean and standard deviation across five simulations.

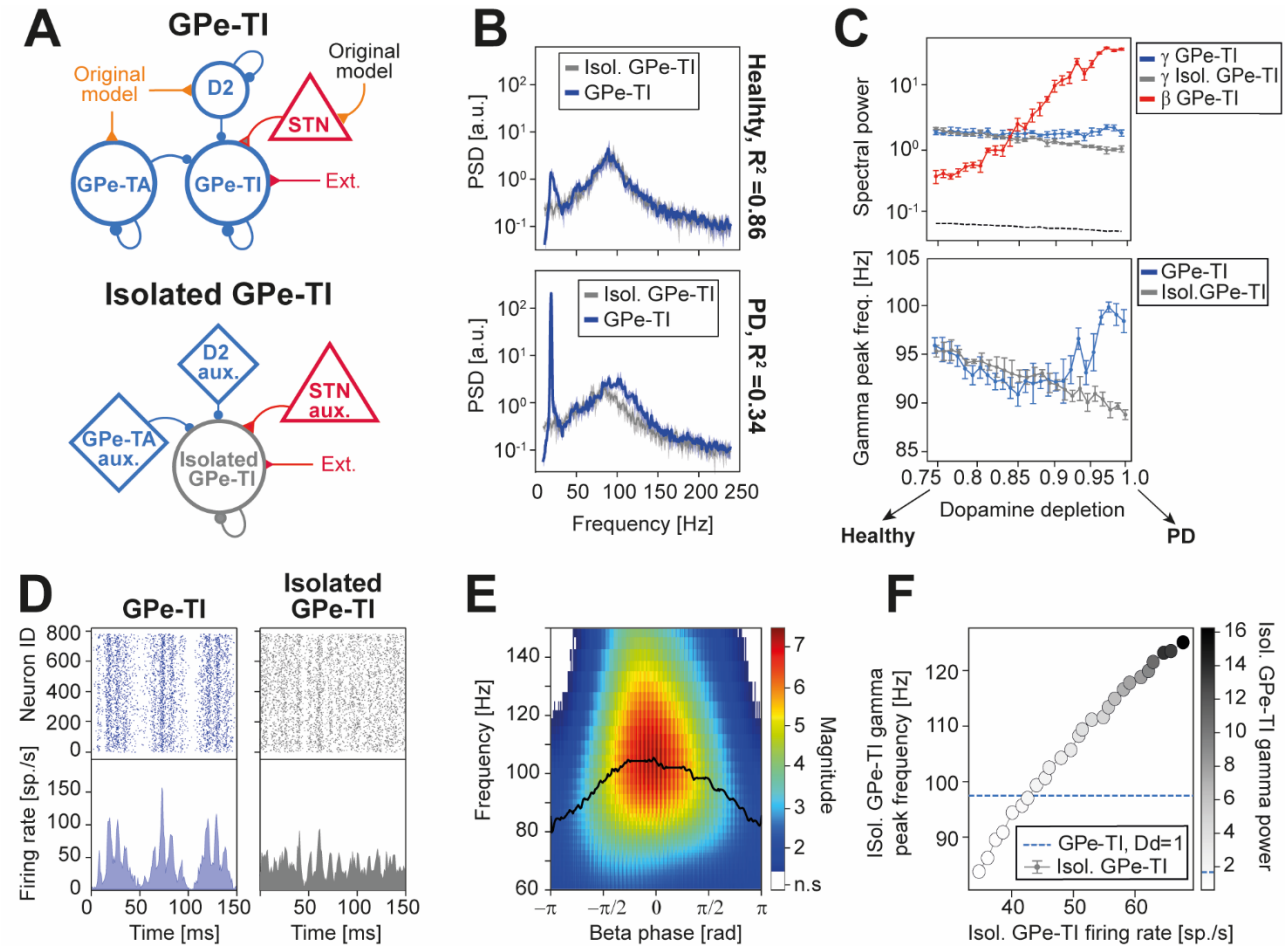

Fig 3. Interplay between beta and gamma oscillations in GPe-TI. (A) Schematic of isolation procedure for GPe-TI: the original nucleus (top) was detached by the rest of the model and the inputs from other nuclei were substituted with Poissonian sources (indicated with "aux." in the bottom panel) with the same firing rates. External drive and self-inhibition were left unchanged. (B) Comparison of the power spectral densities (PSDs) of GPe-TI (blue) and isolated GPe-TI (grey) under healthy (D_d=0.75, bottom) and Parkinsonian conditions (D_d=1, bottom). (C) Spectral power (top) and gamma peak frequency (bottom) of GPe-TI (blue) and isolated GPe-TI (grey) across D_d. The spectral power in the beta band (red) and the threshold for significant oscillations (black dashed lines), computed as the spectral power of a Poissonian process with the same firing rate, are also

reported. Results are reported as mean across five simulations of 5 seconds. (D) Raster plots (top) and activity traces (bottom) of GPe-TI (left) and isolated GPe-TI (right), in Parkinsonian condition. (E) Scalogram of GPe-TI activity averaged across beta cycles, under PD conditions. White regions indicate non-significant oscillations ("n.s."), i.e., those time-frequency regions for which the wavelet amplitudes are below the amplitude of a Poissonian process with the same mean. The black line represents the frequency with the maximum wavelet amplitude for each beta phase (smoothed with a moving average for display purposes). (F) Increase of gamma peak frequency and gamma spectral power with a higher mean firing rate of isolated GPe-TI. The increase of the firing was obtained by increasing the input to the isolated population. For comparison, the original nucleus spectral power and peak frequency are represented with a blue dashed line, for Parkinsonian condition.

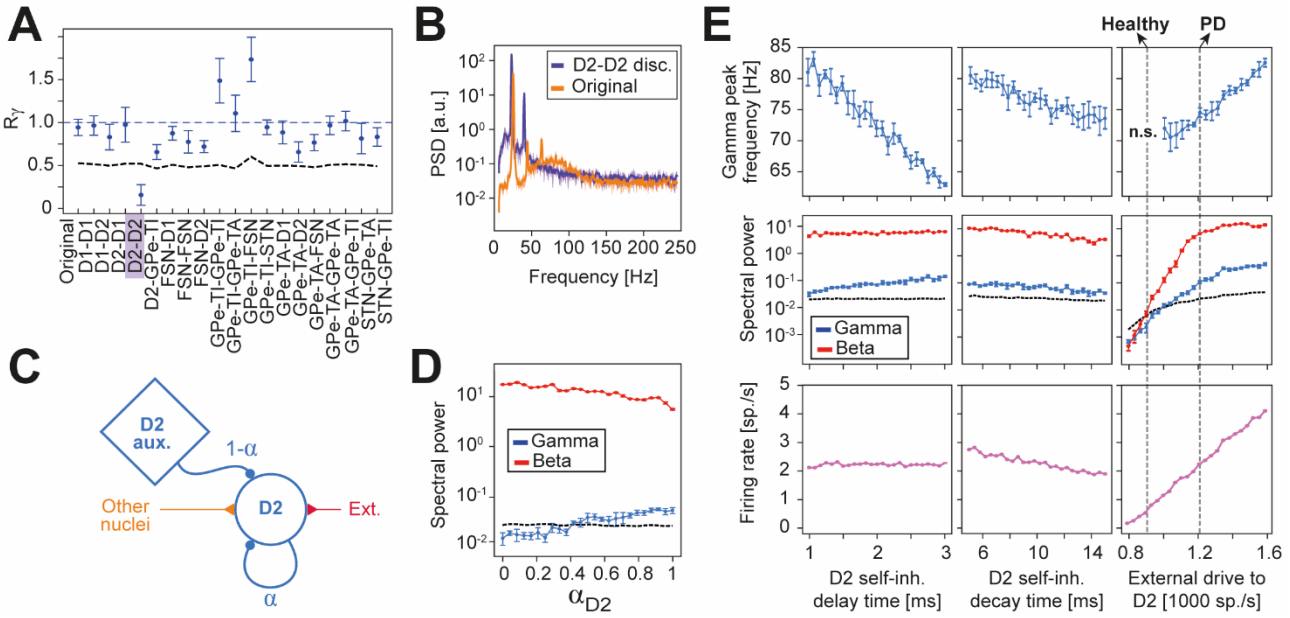

Fig 4. Self-inhibition parameters and external drive modulate D2 oscillations. (A) Ratio Rɣ of D2 gamma power with each synaptic connection silenced compared to the gamma power in the intact full-BG model. The spectral power ratio of D2 activity in the gamma band is shown with ('original') and without each synaptic connection in the full-BG model, with the specific connection removed indicated on the x-axis. For each removed connection, the threshold (black dashed line) for significant oscillations is computed as the ratio between the power of equivalent poisson processes and the gamma power in the fully connected model. The blue dashed line indicates the spectral power ratio for the original fully connected case. (B) Power spectral

834 densities (PSDs) of D2 with (orange) and without (purple) self-inhibitory connection in Parkinsonian condition
835 (D_d=1). (C) Scheme illustrating the procedure for reducing D2 self-inhibition: the weights of self-inhibitory
836 connections were progressively reduced by a multiplicative factor $\alpha \in [0,1]$, and input from an auxiliary
837 Poissonian population (D2 aux.) was added to compensate for the reduction in self-inhibitory input. The
838 afferent activity from other BG nuclei and external inputs were left unchanged. (D) D2 spectral power of beta
839 (red) and gamma (blue) bands across self-inhibition synaptic strength α. The threshold for statistically
840 significant oscillatory power (dashed black line) was determined considering the power spectrum of an
841 equivalent Poisson process with a matched firing rate. (E) Gamma peak frequency (top row), beta (red) and
842 gamma (blue) spectral powers (middle row), and D2 firing rate (bottom row) across D2 self-inhibition delay
843 (left column), D2 self-inhibition decay time (middle column), and the strength of external input to D2 (right
844 column). In the middle row, the threshold for significant oscillation was indicated with a black dashed line.
845 All data was reported as mean and standard deviation across five simulations. In the top row, peak frequencies
846 are not reported for non-significant ("n.s", according to comparison with the power of an equivalent Poissonian
847 process with matched firing rate) gamma oscillations. The grey dashed lines show the value for which the
848 external input match the healthy and pathological conditions.
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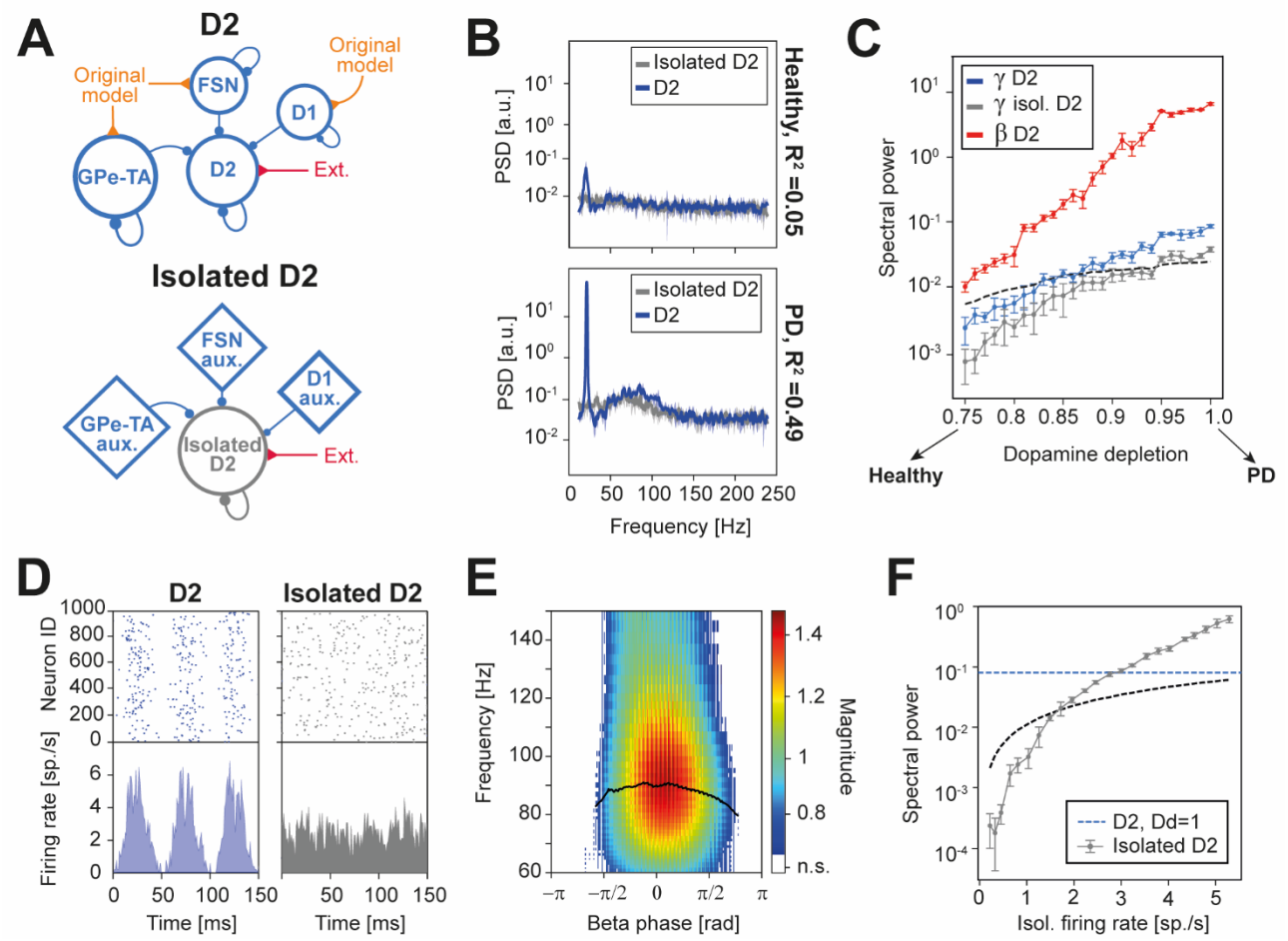


Fig 5. Interplay between beta and gamma oscillations in D2. (A) Schematic of isolation procedure for D2: the original nucleus (top) was detached by the rest of the model and the inputs from other nuclei were substituted with Poissonian sources (indicated with "aux." in the bottom panel) with the same firing rates. External drive and self-inhibition were left unchanged. (B) Comparison of the power spectral densities of D2 (blue) and isolated D2 (grey) under healthy (D_d=0.75, bottom) and Parkinsonian conditions (D_d=1, bottom). (C) Spectral power in the gamma band of D2 (blue) and isolated D2 (grey) across D_d. The spectral power in the beta band (red) and the threshold for significant oscillations (black dashed line), computed as the spectral power of a Poissonian process with the same firing rate, are also reported. Results are reported as mean across five simulations of 5 seconds. (D) Raster plots (top) and activity traces (bottom) of D2 (left) and isolated D2 (right), in parkinsonian condition. (E) Scalogram of D2 activity averaged across beta cycles under Parkinsonian conditions. White regions indicate non-significant oscillations ("n.s."), i.e., those time-frequency regions for which the wavelet amplitudes are below the amplitude of a Poissonian process with the same mean. The black line represents the maximum wavelet amplitude frequency for each beta phase (smoothed with a moving

average for display purposes). (F) Increase of gamma spectral power with a higher mean firing rate of isolated D2. The original nucleus spectral power for the Parkinsonian condition (blue dashed line) and the threshold for significant oscillations (black dashed line) are reported.

Supporting Information

Nucleus	Experimental firing rate [sp./s]	References	Model firing rate [sp./s]
D1	0.5-3	(93,94)	2.41 ± 0.01
D2	0.5-3	(93,94)	0.526 ± 0.004
FSN	10-20	(95,96)	17.66 ± 0.05
GPe-TI	30-60	(70,97)	45.58 ± 0.03
GPe-TA	10-20	(97)	15.46 ± 0.03
STN	10-20	(98,99)	19.44 ± 0.01

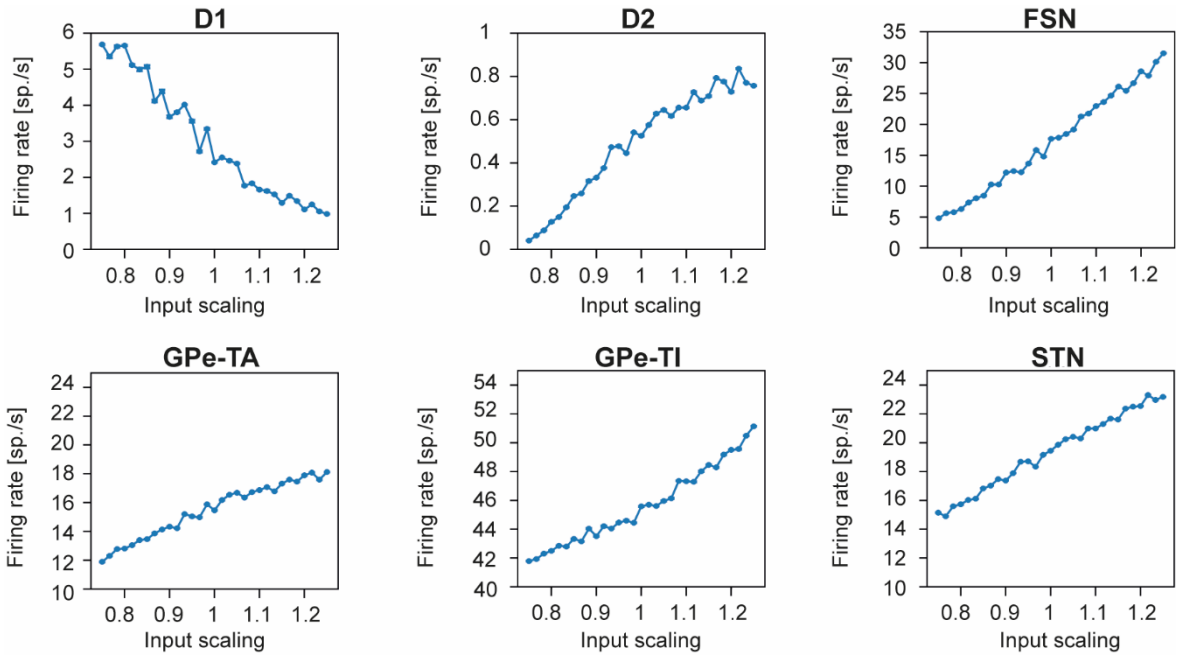
S1 Table. Experimental literature data on basal ganglia nuclei firing rates and corresponding model values under healthy conditions.

Gamma oscillations central	Source nucleus	Human/rats	Reference
~100 Hz	Striatum	Rats	(37)
~70 Hz	Striatum	Rats	(54)
80-100 Hz	Striatum	Rats	(38)
80-110 Hz	Striatum and External Globus	Rats	(39)
80-110 Hz	Subthalamic Nucleus	Rats	(37)
60-90 Hz	Globus Pallidus	Rats	(40)
60-90 Hz	Subthalamic Nucleus	Human	(43)
60-90 Hz	Subthalamic Nucleus	Human	(33)
60-90 Hz	Subthalamic Nucleus	Human	(100)

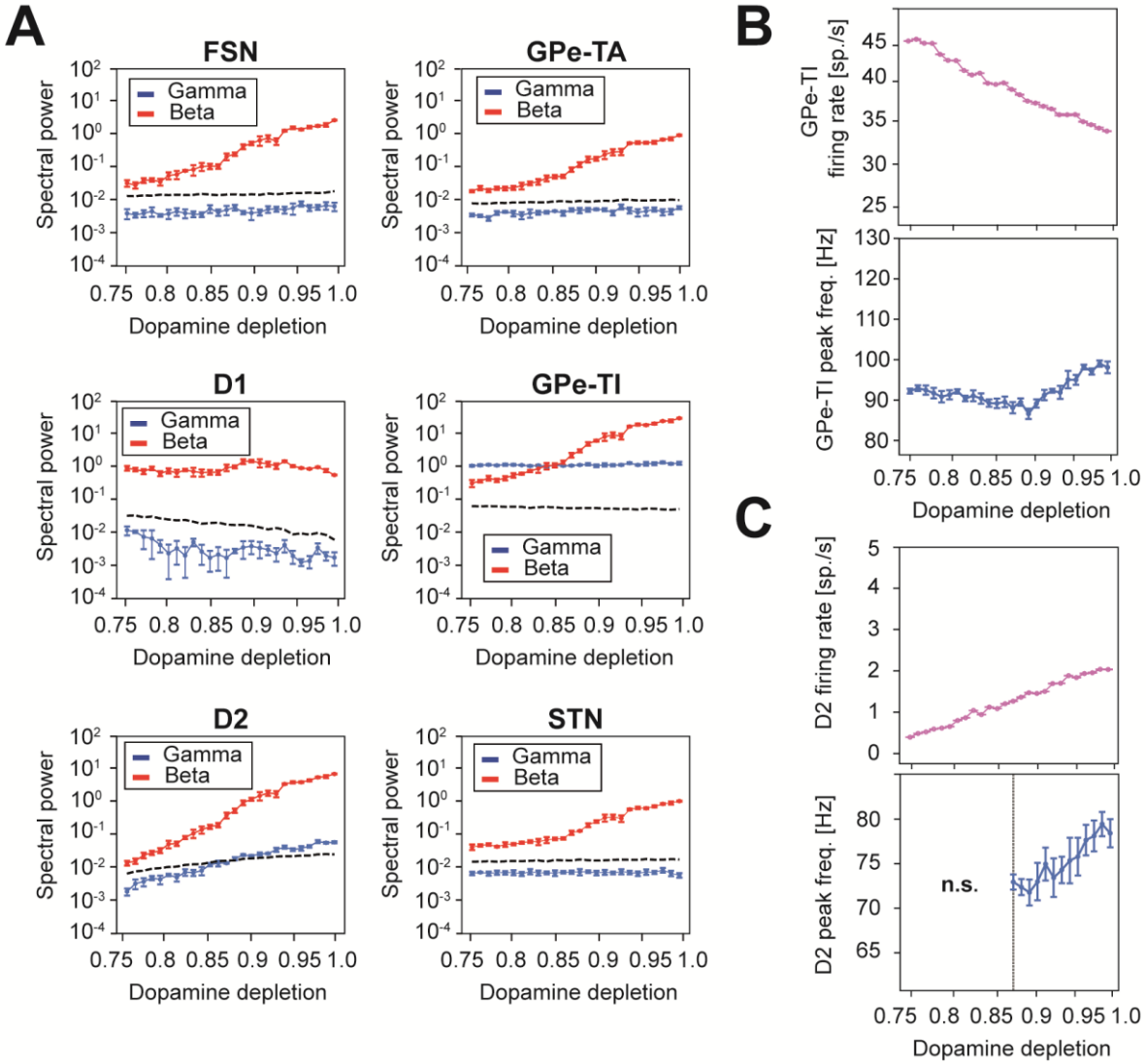
S2 Table. Frequency of gamma oscillations with the basal ganglia in experimental literature.

Parameter	Previous model value	New model value
GPe-TI to GPe-TI synaptic delay	1 ms	1.8 ms
GPe-TI synaptic decay time	5.5 ms	7 ms
GPe-TI to GPe-TI synaptic weight	1.2×10^{-3}	1.1×10^{-3}
GPe-TI external input rate	1530 spikes/s	600 spikes/s
D2 to D2 synaptic weight	0.2×10^{-3}	0.25×10^{-3}
D2 external input rate in PD conditions	1083 spikes/s	1200 spikes/s
D1 external input rate	1127 spike/s	1280 spikes/s

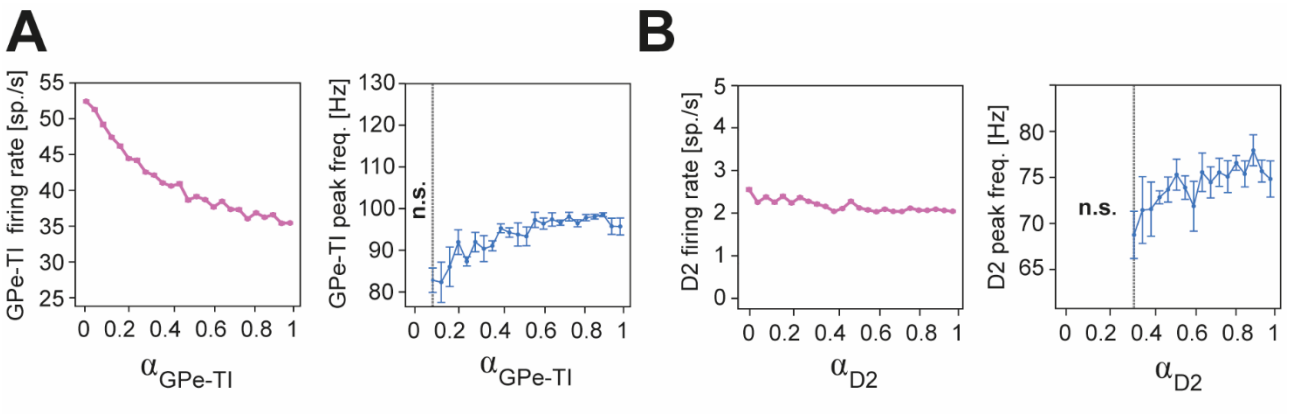
S3 Table. Modified parameters in the updated model.



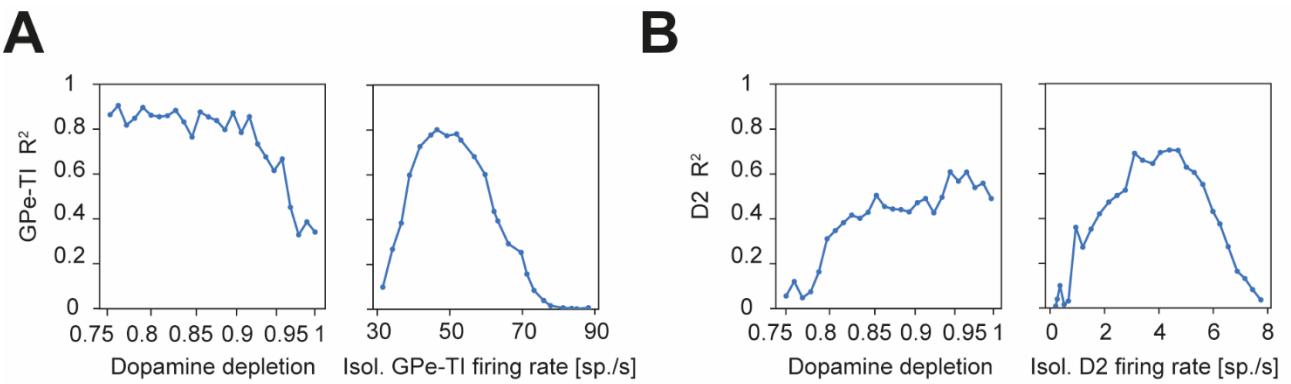
S1 Fig. Impact of external input scaling on network firing properties. Average firing rate of each nucleus in the model under healthy conditions, shown as a function of a multiplicative scaling factor applied to the external input rate. A scaling factor of 1 corresponds to the baseline external input rates reported in Table 3. Results represent the mean firing rate across five simulations, each lasting 5 seconds.



S2 Fig. GPe-TI and D2 are the only nuclei displaying gamma activity. A) Spectral power in the gamma (blue) and beta (red) bands for each basal ganglia nucleus of the model, across different dopamine depletion levels. The threshold for significant gamma oscillations (black dashed line) is computed as the gamma spectral power of a Poissonian process with the same firing rate. Results are presented as mean and standard deviation across five simulations of 5 seconds. B) GPe-TI firing rate (top) and gamma peak frequency (bottom) with different levels of dopamine depletions. C) D2 firing rate (top) and gamma peak frequency (bottom) with different levels of dopamine depletions. Peak frequencies are not reported for non-significant ("n.s", according to comparison with the power of an equivalent Poissonian process with matched firing rate) gamma oscillations.

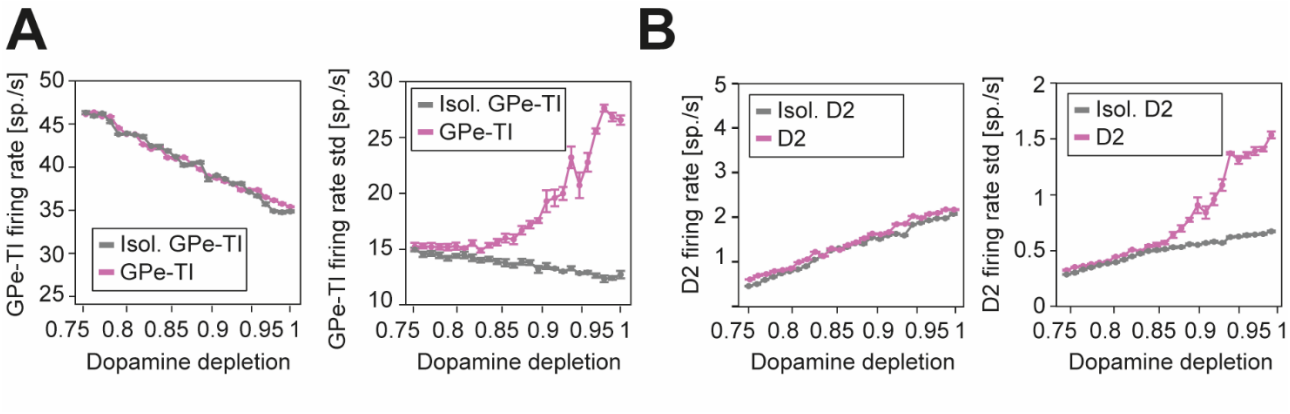


S3 Fig. Effects of self-inhibition disconnection on firing rates and gamma peak frequencies of GPe-TI and D2. A) GPe-TI gamma peak frequency (left) and firing rate (right) across different levels of self-inhibition synaptic strength α_{GPe-TI}. Peak frequencies are not reported for non-significant ("n.s", according to comparison with the power of an equivalent Poissonian process with matched firing rate) gamma oscillations. Results are presented as mean and standard deviation across five simulations of 5 seconds. B) D2 gamma peak frequency (left) and firing rate (right) across different levels of self-inhibition strength α_{D2}. Peak frequencies are not reported for non-significant ("n.s", according to comparison with the power of an equivalent Poissonian process with matched firing rate) gamma oscillations. Results are presented as mean and standard deviation across five simulations of 5 seconds.



S4 Fig. Similarity between the isolated and original power spectral densities across dopamine depletion levels and firing rates of the isolated nuclei. A) left, coefficient of determination R^2 between the power spectral densities (considering the range [50-150] Hz) of the isolated GPe-TI and the full BG GPe-TI across dopamine depletion levels; right, R^2 between the power spectral densities of GPe-TI at D_d=1 and isolated GPe-TI, varying the firing rate of the latter and considering the range [50-150] Hz. B) left, R^2 between the power

spectral densities (considering the range [50-150] Hz) of D2 and isolated D2, across different levels of dopamine depletion; right, R^2 between the power spectral densities of D2 at D_d=1 and isolated D2, varying the firing rate of the latter and considering the range [50-150] Hz.



S5 Fig. Isolated versions of D2 and GPe-TI present the same average firing activities of the original nuclei, but smaller fluctuations.

A) Mean firing rate (left) and its simulation standard deviation (right) of GPe-TI (pink) and isolated GPe-TI (grey) across dopamine depletion levels (D_d). The isolated GPe-TI population represents an equivalent of GPe-TI, but with inputs from other nuclei replaced by Poissonian sources with matching firing rates. Results are presented as the mean and standard deviation across five simulations, each lasting 5 seconds. B) Mean firing rate (left) and its simulation standard deviation (right) of D2 (pink) and isolated D2 (grey) across dopamine depletion levels (D_d). The isolated D2 population represents an equivalent of D2, but with inputs from other nuclei replaced by Poissonian sources with matching firing rates. Results are presented as the mean and standard deviation across five simulations, each lasting 5 seconds.

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